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Reviewer: Anne Corrigan

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Application No: 10534292 Version No: 1.1

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	10534292 2005-05-09
	PCT/BE2003/000190 2003-11-07
	US 60/425,073 2002-11-08
	US 60/425,063 2002-11-08
	EP 03447005.4 2003-01-10
<150> <151>	PCT/EP03/06581 2003-06-23
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Val Met Ala	Trp Phe	Arg Gln	Ala I	Pro Gly	Lys	Gln	Arg 45	Asp	Phe	Val		
Ala Tyr Ile	e Thr Ser	Ala Val 55	Asn '	Thr Asp	Tyr	Ala 60	Asp	Phe	Val	Lys		
Gly Arg Phe	e Thr Ile	Ser Arg 70	Asp A	Asn Ala	Gln 75	Asn	Met	Val	His	Leu 80		
Gln Met Asr	Ser Leu 85	Lys Pro	Glu Z	Asp Thr 90	Ala	Val	Tyr	Tyr	Cys 95	Asn		
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Ser Leu Ala Trp Phe Arg Gln Ala Pro Gly Lys Glu Arg Asp Phe Val 40 45

35

Ala Ala Leu Ser Leu Thr Thr Tyr Tyr Ala Asp Ser Val Lys Gly Arg 50 55 60 Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn Thr Val Tyr Leu Gln Met 70 75 80 Asn Ser Leu Lys Pro Asp Asp Thr Ala Asp Tyr Phe Cys Ala Thr Ala 85 90 Arg Thr Arg Thr Asp Tyr Ala Pro Leu Ser Ala Ala Ser Thr Tyr 105 110 Asp Ala Trp Gly Gln Gly Thr Gln Val Thr Val Ser Leu 115 120 125 <210> 4 <211> 124 <212> PRT <213> Lama glama <400> 4 Gln Val Gln Leu Gln Glu Ser Gly Gly Gly Leu Val Gln Ala Gly Gly 1 5 10 15 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Arg Ser Ser Arg Tyr Tyr 20 25 30 Ala Met Gly Trp Phe Arg Gln Gly Pro Gly Lys Glu Arg Glu Phe Val 35 40 45 Ala Ala Val Asn Trp Asn Gly Asp Ser Thr Tyr Tyr Ala Asp Ser Val 50 55 60 Lys Gly Arg Phe Thr Ile Ser Arg Gly Asn Ala Glu Asn Thr Ala Tyr 70 75 65 Leu Gln Met Asn Ser Leu Val Pro Glu Asp Thr Ala Val Tyr Tyr Cys

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Val Met Ala Trp Phe Arg Gln Thr Pro Gly Lys Glu Arg Glu Phe Val 35 40 45

Gly Ala Ile Asp Trp Ser Gly Arg Arg Ile Thr Tyr Thr Asp Ser Val50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn Thr Val Tyr 65 70 75 80

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Val Met Ala Trp Phe Arg Gln Thr Pro Gly Lys Glu Arg Glu Phe Val
35 40 45

Gly Ala Ile Asp Trp Ser Gly Arg Arg Ile Thr Tyr Thr Asp Ser Val
50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn Thr Val Tyr 65 70 75 80

Leu Gln Met Asn Thr Leu Lys Pro Glu Asp Thr Ala Val Tyr Tyr Cys 85 90 95

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Ala Ser Ile Ser Ser Ser Gly Ile Ser Thr Tyr Tyr Ala Asp Ser Val 50

Lys Gly Arg Phe Thr Ile Ser Arg Asp Ile Ala Lys Asn Thr Val Tyr 75

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70

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Leu Gln Met Asn Ser Leu Lys Pro Glu Asp Thr Ala Val Tyr Tyr Cys
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Lys Gly Arg Phe Thr Ile Ser Asn Asp Lys Val Lys Asn Thr Val Tyr 65 70 75 80

Leu Gln Met Asn Ser Leu Lys Pro Glu Asp Thr Ala Val Tyr Phe Cys
85 90 95

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Ala Met Gly Trp Phe Arg Gln Ala Pro Gly Lys Glu Arg Glu Phe Val 35 40 45

Ala Ala Val Ser Tyr Ser Gly Ser Tyr Tyr Ala Asp Ser Val Lys Gly 50 55 60

Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn Thr Val Tyr Leu Gln 65 70 75 80

Met Ala Ser Leu Lys Pro Glu Asp Thr Ala Val Tyr Tyr Cys Ala Ala 85 90 95

Arg Asn Arg Gly Tyr Ser Thr Tyr Ala Gly Val Tyr Asp Tyr Trp Gly 100 105 110

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     35 40 45
Ala Ser Ile Thr Trp Ile Gly Gly Gly Thr Tyr Tyr Ala Asp Ser Val
      55 60
Lys Gly Arg Phe Thr Ile Ser Arg Asp His Ala Gly Asn Thr Val Tyr
         70
                      75 80
Leu Gln Met Asn Thr Leu Lys Pro Asp Asp Thr Ala Val Tyr Tyr Cys
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Ala Leu Asp Arg Arg Ser Ser Thr Tyr Tyr Leu Met Lys Gly Glu Tyr
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Arg Glu Phe Val Ala Arg Ile Tyr Trp Ser Ser Gly Asn Thr Tyr Tyr 50 55 60

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Ala Asp Ser Val Lys Gly Arg Phe Ala Ile Ser Arg Asp Ile Ala Lys 7.0 Asn Thr Val Asp Leu Thr Met Asn Asn Leu Glu Pro Glu Asp Thr Ala 85 90 Val Tyr Tyr Cys Ala Ala Arg Asp Gly Ile Pro Thr Ser Arg Ser Val 100 105 110 Glu Ser Tyr Asn Tyr Trp Gly Gln Gly Thr Gln Val Thr Val Ser Ser 115 120 Ala Ala Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Asn Gly Ala 135 140 Ala His His His His His 145 150 <210> 13 <211> 124 <212> PRT <213> Lama glama <400> 13 Gln Val Gln Leu Gln Asp Ser Gly Gly Gly Leu Val Gln Ala Gly Gly 10 Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Gly Thr Phe Ser Ser Ile 20 25 Ile Met Ala Trp Phe Arg Gln Ala Pro Gly Lys Glu Arg Glu Phe Val 35 40 45 Gly Ala Val Ser Trp Ser Gly Gly Thr Thr Val Tyr Ala Asp Ser Val

Leu Gly Arg Phe Glu Ile Ser Arg Asp Ser Ala Arg Lys Ser Val Tyr

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65

75

Leu Gln Met Asn Ser Leu Lys Pro Glu Asp Thr Ala Val Tyr Tyr Cys 85 90 95

Ala Ala Arg Pro Tyr Gln Lys Tyr Asn Trp Ala Ser Ala Ser Tyr Asn 100 105 110

Val Trp Gly Gln Gly Thr Gln Val Thr Val Ser Ser \$115\$

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Ile Met Ala Trp Phe Arg Gln Ala Pro Gly Lys Glu Arg Glu Phe Val 35 40 45

Gly Ala Val Ser Trp Ser Gly Gly Thr Thr Val Tyr Ala Asp Ser Val
50 55 60

Leu Gly Arg Phe Glu Ile Ser Arg Asp Ser Ala Arg Lys Ser Val Tyr 65 70 75 80

Leu Gln Met Asn Ser Leu Lys Pro Glu Asp Thr Ala Val Tyr Tyr Cys 85 90 95

Ala Ala Arg Pro Tyr Gln Lys Tyr Asn Trp Ala Ser Ala Ser Tyr Asn 100 105 110

Val Trp Gly Gln Gly Thr Gln Val Thr Val Ser Ser Glu Pro Lys Thr 115 120 125

Pro Lys Pro Gln Pro Ala Ala Gln Val Gln Leu Gln Asp Ser Gly

130 135 140

Ser Gly Gly Thr Phe Ser Ser Ile Ile Met Ala Trp Phe Arg Gln Ala 165 170 175

Pro Gly Lys Glu Arg Glu Phe Val Gly Ala Val Ser Trp Ser Gly Gly 180 185 190

Thr Thr Val Tyr Ala Asp Ser Val Leu Gly Arg Phe Glu Ile Ser Arg 195 200 205

Asp Ser Ala Arg Lys Ser Val Tyr Leu Gln Met Asn Ser Leu Lys Pro 210 215 220

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